



Note to the user

This software has been written to analyse data of the SPI telescope onboard INTEGRAL. Particular care has been taken in making the software user friendly and well documented. If you appreciated this effort, and if this software and User Manual were useful for your scientific work, the author would appreciate a corresponding acknowledgment in your published work.

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1 Introduction

The executable spi_tm2fits converts INTEGRAL or SPI telemetry in the format of the CESR GSE aquisition system (PktAll) into a ISDC compliant data structure of ISDC level RAW. Each run of spi_tm2fits produces a single science windows, hence to cut a data stream into science window it has to be applied successively to the input data with different input packet limitations (use the task spi_tmscan to automatically devide a data stream into science windows).

spi_tm2fits comes with a number of data structure templates that may either replace or supplement existing ISDC templates. Templates that are replaced are ISDC templates that do not reflect the actual status of the data base. Templates that are supplemented concern data that is normally not analysed by the ISDC system, such as CNES GSE packets or sub-assembly memory dumps.

spi_tm2fits has been written in the ANSI C++ language, and requires the libraries spi_psdlib and spi_toolslib. The actual version is compliant with the ISDC support platform 4.1.

2 Getting started

Before installing spi_tm2fits, make sure that the ISDC support platform 4.1 or higher is installed on your system, and that the libraries spi_psdlib and spi_toolslib are also available (please consult the WWW site http://www.cesr.fr/~jurgen/isdc if these libraries are not available on your system).

After downloading the spi_tm2fits.tar.gz file, step into a directory that should hold the distribution, move the spi_tm2fits.tar.gz file into this directory and type:

```
$ gunzip spi_tm2fits.tar.gz
$ tar xvf spi_tm2fits.tar
```

The first command uncompresses the distribution file, the second unpacks the files.

Before configuration, the distribution needs to be reset to a clean state. To do this, type

```
$ make distclean
```

Then, configure the distribution. It is assumed here that you have previously installed the ISDC support platform, thus you should type

```
$ ~/bin/ac_stuff/configure
```

Finally, build the distribution by typing

```
$ make global_install
```

3 Parameter file

```
#
              Centre d'Etude Spatiale des Rayonnements
                                                                #
#
                    (in collaboration with ISDC)
                                                                #
#
#
                         SPI Pre Processing
#
# File: spi_tm2fits.par
# Version: 1.5.0
                                                                #
# Component: pp
#
# Author:
            Juergen Knoedlseder
#
            knodlseder@cesr.fr
#
            CESR
#
# Purpose: Parameter file of the SPI pre-processing executable
# History: 1.5.0 18-Jul-2002
# Path name definition
#==========
                                   "/x02/tm",,, "TM Base Path"
tmpath, s, h,
scwbase, s, h, "/villa-1/jurgen/arc/tst_tm/scw",,, "Science Window Base Path"
cfgbase, s, h, "/users/jurgen/isdc/templates",,, "Template Base Path"
# TM input definition
#=========
                        515,,, "First TM file number"
nfirst, i, ql,
                       515,,, "Last TM file number"
nlast, i, ql,
pktmin, i, ql, 400174,1,10000000, "First packet to scan in first TM file"
pktmax, i, ql, 447092,1,10000000, "Last packet to scan in last TM file"
header, i, h, 0,,, "Header size (in Bytes)"
# Science window definition
revid, s, ql, "9000",,, "Revolution ID"

pointid, s, ql, "0080",,, "Pointing ID (minimum is 0000)"

subid. s, ql, "001",,, "Subdivision ID (minimum is 001)"

"Goionce Window Type (0=pointing,
scwtype, s, ql, "2",0|1|2,, "Science Window Type (0=pointing, 1=slew, 2=other)"
version, s, ql, "001",,,
                             "Version number"
previd, s, ql, "00000000000",,,"Previous Science Window ID"
# Task parameters
#========
method, s, ql, "PROCESS", SCAN | PROCESS,, "Working method (SCAN / PROCESS)"
bcpid, s, ql, "00000000",,, "Broadcast packet ID"
                                "Reason for Science Window ending"
swbound, s, ql, "UNDEFINED",,,
```

The parameters have the following meanings:

- tmpath specified the directory which holds the PktAll telemetry files in the CESR GSE format (in this format, telemetry frames are stored as successive blocks of 440 Bytes in a single binary file).
- scwbase specifies the Science window base path into which the science window data structures should be written.
- cfgbase specifies the absolute directory in which the science window group configuration files will be found. Normally, the directory where the data structure templates are found should be specified.
- nfirst specifies the file number of the first PktAll telemetry file that should be processed.
- nlast specifies the file number of the last PktAll telemetry file that should be processed.
- pktmin specifies the first packet in the first telemetry binary file (specified by the parameter nfirst) that should be processed (the first packet has the packet number 1). A packet is defined as a 440 Bytes block in the telemetry binary file. If pktmin = 1, pre-processing starts from the beginning of the telemetry file.
- pktmax specifies the last packet in the last telemetry binary file (specified by the parameter nlast) that should be processed (packets start from number 1). A packet is defined as a 440 Bytes block in the telemetry binary file. If pktmax is larger than the number of available packets, the telemetry file is analysed until the end of the file is reached.
- header specifies the number of Bytes in the telemetry binary file header. Earlier versions of the PktAll files had a header attached. The actual version do not have a header anymore, hence as default, 0 should be specified.
- revid specifies the revolution identifier of the science window that should be generated. This parameter has to be a 4 character string.
- pointid specifies the pointing identifier of the science window that should be specified. This parameter has to be a 4 character string. The minimum pointing identifier is 0000.
- subid specifies the subdivision identifier of the science window that should be specified. This parameter has to be a 3 character string. The minimum subdivision identifier is 001.
- scwtype specifies the science window type. The following types are allowed: 0 is a pointing, 1 is a slew, and 2 is any other science window type.
- version specifies the science window version number. This parameter has to be a 3 character string.
- previd specifies the previous science window identifier (without version number). This identifier will be stored in the science window grouping table header, and allows for linking of the science windows.
- method specifies the telemetry processing method. If SCAN is specified, the telemetry file is only scanned and statistics about the contained data types will be dumped into the RIL log file. No RAW data will be written. If PROCESS is specified, the telemetry file will first be scanned to determine the DAL table size of the resulting RAW data structures, and in a second pass, the telemetry will be pre-processed and written into the data structures.

- bcpid specifies the Broadcast packet identifier for the telemetry, to be stored in the headers of the RAW data structures.
- swbound specifies the reason for science window ending. Possible reasons are SLEW, POINTING, TIMEOUT, and UNDEFINED.
- clobber ISDC standard parameter (see Common User Manual).
- mode ISDC standard parameter (see Common User Manual).

4 Interface definition

spi_tm2fits creates a science window group of ISDC level RAW from the input telemetry files that is compliant with the ISDC data structure defintions. The following HDUs are filled by the executable in the public eng branch of the science window structure:

Filename	HDU	Content
eng/raw/intl_stsp.fits	INTL-STSP-SRW	Spacecraft time packet
eng/raw/intl_raw_hk.fits	INTL-PLMHRW	Spacecraft PLM packet
G	INTL-CDMU-HRW	Spacecraft CDMU packet
	INTL-AOCS-HRW	Spacecraft AOCS packet
	INTL-SVM1-HRW	Service module 1 packets (only SPI parameters)
	INTL-SVM2-HRW	Service module 1 packets (only SPI parameters)
eng/raw/spi_raw_acs.fits	SPIFEE0-HRW	ACS group 0 count rates
•	SPIFEE1-HRW	ACS group 1 count rates
	SPIFEE2-HRW	ACS group 2 count rates
	SPIFEE3-HRW	ACS group 3 count rates
	SPIFEE4-HRW	ACS group 4 count rates
	SPIFEE5-HRW	ACS group 5 count rates
	SPIFEE6-HRW	ACS group 6 count rates
	SPIFEE7-HRW	ACS group 7 count rates
	SPIFEE8-HRW	ACS group 8 count rates
	SPIFEE9-HRW	ACS group 9 count rates
	SPIFEEA-HRW	ACS group 10 count rates
	SPIFEEB-HRW	ACS group 11 count rates
	SPIOACS-HRW	ACS overall count rates
eng/raw/spi_raw_hk.fits	SPIDPEHRW	SPI DPE HK (1 second)
	SPI001HRW	SPI HK (8 seconds)
	SPI008HRW	SPI HK (64 seconds)
	SPI080A-HRW	SPI HK (640 seconds - a)
	SPI080B-HRW	SPI HK (640 seconds - b)
	SPI480A-HRW	SPI HK (3840 seconds - a)
	SPI480B-HRW	SPI HK (3840 seconds - b)
	SPI480C-HRW	SPI HK (3840 seconds - c)
	SPI480D-HRW	SPI HK (3840 seconds - d)
eng/raw/spi_raw_dhk.fits	SPIDHKO-CRW	DIAG housekeeping packet 0
	SPIDHK1-CRW	DIAG housekeeping packet 1
	SPIDHK2-CRW	DIAG housekeeping packet 2
	SPIDHK3-CRW	DIAG housekeeping packet 3
	SPIDHK4-CRW	DIAG housekeeping packet 4
	SPIDHK5-CRW	DIAG housekeeping packet 5
	SPIDHK6-CRW	DIAG housekeeping packet 6
	SPIDHK7-CRW	DIAG housekeeping packet 7
	SPIDHK8-CRW	DIAG housekeeping packet 8
	SPIDHK9-CRW	DIAG housekeeping packet 9
	SPIDHKA-CRW	DIAG housekeeping packet 10
	SPIDHKB-CRW	DIAG housekeeping packet 11

$\operatorname{Filename}$	HDU	Content
eng/raw/spi_acs_calib.fits	SPIACSCRW	ACS calibration data
eng/raw/spi_raw_dump.fits	SPIDFMD-HRW	DFEE memory dumps (non ISDC)
eng/iaw/spilawlamp.iios	SPIASMD-HRW	ACS memory dump (non ISDC)
	SPIPDMD-HRW	PSD memory dump (non ISDC)
//		- 1
eng/raw/spi_raw_cnes.fits	CNES-PDUHRW	CNES PDU data (non ISDC)
	CNES-RTUHRW	CNES RTU data (non ISDC)
eng/raw/spi_raw_block.fits	SPIOBLK-HRW	Photon data block structure (OPER)
	SPIEBLK-HRW	Photon data block structure (EMER)
	SPICBLK-HRW	Photon data block structure (CALB)
	SPIDBLK-HRW	Photon data block structure (DIAG)
eng/raw/spi_raw_schk.fits	SPISHK1-TMP	Science HK (temporary 1)
	SPISHK2-TMP	Science HK (temporary 2)
	SPISHK3-TMP	Science HK (temporary 3)
	SPISHK4-TMP	Science HK (temporary 4)
	SPISHK5-TMP	Science HK (temporary 5)

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The following HDUs are filled by spi_tm2fits in the private spi branch of the science window structure:

Filename	HDU	Content
spi/raw/spi_raw_oper.fits	SPIOSGL-RAW	Single events (OPER)
	SPIOPSD-RAW	PSD events (OPER)
	SPIOCRV-RAW	PSD curves (OPER)
	SPIOME2-RAW	Double events (OPÉR)
	SPIOME3-RAW	Tripple events (OPER)
	SPIOME4-RAW	Quadruple events (OPER)
	SPIOME5-RAW	Quintuple events (OPER)
	SPIOMEH-RAW	Multiple events (OPER)
	SPIOMP3-RAW	Tripple with PSD events (OPER)
	SPIOMP4-RAW	Quadruple with PSD events (OPER)
	SPIOMP5-RAW	Quintuple with PSD events (OPER)
	SPIOMP6-RAW	Setuple with PSD events (OPER)
	SPIOMPH-RAW	Multiple with PSD events (OPER)
	SPIOPPS-RAW	Pure PSD events (OPER)
spi/raw/spi_raw_emer.fits	SPIEME2-RAW	Double events (EMER)
	SPIEME3-RAW	Tripple events (EMER)
	SPIEME4-RAW	Quadruple events (EMER)
	SPIEME5-RAW	Quintuple events (EMER)
	SPIEMEH-RAW	Multiple events (EMER)
	SPIEMP3-RAW	Tripple with PSD events (EMER)
	SPIEMP4-RAW	Quadruple with PSD events (EMER)
	SPIEMP5-RAW	Quintuple with PSD events (EMER)
	SPIEMP6-RAW	Setuple with PSD events (EMER)
	SPIEMPH-RAW	Multiple with PSD events (EMER)
	SPIEPPS-RAW	Pure PSD events (EMER)
spi/raw/spi_raw_calb.fits	SPICCRV-RAW	PSD curves (CALB)
spi/raw/spi_raw_diag.fits	SPIDSGL-RAW	Single events (DIAG)
	SPIDPSD-RAW	PSD events (DIAG)
	SPIDCRV-RAW	PSD curves (DIAG)
	SPIDME2-RAW	Double events (DIAG)
	SPIDME3-RAW	Tripple events (DIAG)
	SPIDME4-RAW	Quadruple events (DIAG)
	SPIDME5-RAW	Quintuple events (DIAG)
	SPIDMEH-RAW	Multiple events (DIAG)
	SPIDMP3-RAW	Tripple with PSD events (DIAG)
	SPIDMP4-RAW	Quadruple with PSD events (DIAG)
	SPIDMP5-RAW	Quintuple with PSD events (DIAG)
	SPIDMP6-RAW	Setuple with PSD events (DIAG)
	SPIDMPH-RAW	Multiple with PSD events (DIAG)
	SPIDPPS-RAW	Pure PSD events (DIAG)

Additional data structure may be craeted by spi_tm2fits, yet they will not be filled with information.

The following header keywords of the science window group are set by spi_tm2fits:

Keyword	Value	Content
INSTRUME	SPI	INTEGRAL Instrument
ERTFIRST		Earth received time start of telemetry
ERTLAST		Earth received time stop of telemetry
SWBOUND		Reason for Science Window ending
PCKSTART		First TM packet
PCKEND		Last TM packet

In addition, the following keywords are added to the RAW data structure headers:

Keyword	Value	Content
ERTFIRST		Earth received time start of telemetry
ERTLAST		Earth received time stop of telemetry
PCKSTART		First TM packet
PCKEND		Last TM packet
SSC_BEG		First SSC in data
$\mathtt{SSC}_{\mathtt{END}}$		Last SSC in data
SSC_GAP		Number of SSC gaps in data

All photon data structures will also have the additional keywords:

Keyword	Value	Content
LOBT_SBK		Local OBT start of block
LOBT_EBK		Local OBT end of block
${\tt BLCK_NUM}$		Number of 125 ms patterns in data
BLCK_GAP		Number of 125 ms pattern gaps in data

5 Error codes

The following error codes may be returned by spi_tm2fits:

```
SPI_TM2FITS_ERROR_BASE -10000 // Error base
SPI_TM2FITS_ERROR_MEM_ALLOC -10000 // Memory allocation error
SPI_TM2FITS_ERROR_EOF -10001 // End of file reached
SPI_TM2FITS_ERROR_FILE_ERROR -10002 // TM file error
SPI_TM2FITS_ERROR_INVALID_BUFFER -10003 // Invalid buffer
```